APPLICANT(S): LEVY, Andrew

10/748,177

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LISTING OF THE PENDING CLAIMS

Below are set forth the currently pending claims.

(Currently amended) A method of determining a potential of a diabetic patient to benefit from vitamin E anti-exidant therapy for treatment of a cardiovascular complication comprising cardiovascular death or myocardial infarction, the method comprising determining a haptoglobin phenotype of the diabetic patient and thereby determining the potential of the diabetic patient to benefit from said vitamin E anti-oxidant therapy, wherein said benefit from said vitamin E anti exident therapy to a patient having a haptoglobin 2-2 phenotype is greater compared to patients having haptoglobin 1-2 phenotype or haptoglobin 1-1 phenotypes.

2. to 5. (Cancelled)

- (Withdrawn) The method of claim 1, wherein said determining said haptoglobin phenotype is effected by determining a haptoglobin genotype of the diabetic patient.
- (Withdrawn) The method of claim 6, wherein said step of determining 7. said haptoglobin genotype of the diabetic patient is effected by a method selected from the group consisting of a signal amplification method, a direct detection method and detection of at least one sequence change.
- 8. (Withdrawn) The method of claim 7, wherein said signal amplification method amplifies a molecule selected from the group consisting of a DNA molecule and an RNA molecule.
- 9. (Withdrawn) The method of claim 7, wherein said signal amplification method is selected from the group consisting of PCR, LCR (LAR), Self-Sustained Synthetic Reaction (3SR/NASBA) and Q-Beta (QB) Replicase reaction.
- 10. (Withdrawn) The method of claim 7, wherein said direct detection method is selected from the group consisting of a cycling probe reaction (CPR) and a branched DNA analysis.

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- 11. (Withdrawn) The method of claim 7, wherein said detection of at least one sequence change employs a method selected from the group consisting of restriction fragment length polymorphism (RFLP analysis), allele specific oligonucleotide (ASO) analysis, Denaturing/Temperature Gradient Gel Electrophoresis (DGGE/TGGE), Single-Strand Conformation Polymorphism (SSCP) analysis and Dideoxy fingerprinting (ddF).
- 12. (Original) The method of claim 1, wherein said determining said haptoglobin phenotype is effected by directly determining the haptoglobin phenotype of the diabetic patient.
- 13. (Original) The method of claim 12, wherein step of determining said haptoglobin phenotype is effected by an immunological detection method.
- 14. (Original) The method of claim 13, wherein said immunological detection method is selected from the group consisting of a radio-immunoassay (RIA), an enzyme linked immunosorbent assay (ELISA), a western blot, an immunohistochemical analysis, and fluorescence activated cell sorting (FACS).
- 15. (Currently amended) A method of determining the importance of reducing oxidative stress by administering vitamin E in a diabetic patient so as to prevent a diabetes-associated cardiovascular complication comprising cardiovascular death or myocardial infarction, the method comprising the step of determining a haptoglobin phenotype of the diabetic patient, thereby determining the importance of reducing the oxidative stress by administering vitamin E in the specific diabetic patient, wherein said importance of reducing oxidative stress by administering vitamin E is greater in a patient having a haptoglobin 2-2 phenotype compared to patients having haptoglobin 1-2 phenotype or haptoglobin 1-1 phenotypes.

16. to 19. (Cancelled)

- 20. (Withdrawn) The method of claim 15, wherein said step of determining said haptoglobin phenotype is effected by determining a haptoglobin genotype of the diabetic patient.
- 21. (Withdrawn) The method of claim 15, wherein said step of determining said haptoglobin genotype of the diabetic patient is effected by a method

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selected from the group consisting of a signal amplification method, a direct detection method and detection of at least one sequence change.

- 22. (Withdrawn) The method of claim 21, wherein said signal amplification method amplifies a molecule selected from the group consisting of a DNA molecule and an RNA molecule.
- 23. (Withdrawn) The method of claim 21, wherein said signal amplification method is selected from the group consisting of PCR, LCR (LAR), Self-Sustained Synthetic Reaction (3SR/NASBA) and Q-Beta (Qβ) Replicase reaction.
- 24. (Withdrawn) The method of claim 21, wherein said direct detection method is selected from the group consisting of a cycling probe reaction (CPR) and a branched DNA analysis.
- 25. (Withdrawn) The method of claim 21, wherein said detection of at least one sequence change employs a method selected from the group consisting of restriction fragment length polymorphism (RFLP analysis), allele specific oligonucleotide (ASO) analysis, Denaturing/Temperature Gradient Gel Electrophoresis (DGGE/TGGE), Single-Strand Conformation Polymorphism (SSCP) analysis and Dideoxy fingerprinting (ddF).
- 26. (Original) The method of claim 15, wherein said step of determining said haptoglobin phenotype is effected by directly determining the haptoglobin phenotype of the diabetic patient.
- 27. (Original) The method of claim 26, wherein said step of determining said haptoglobin phenotype is effected by an immunological detection method.
- 28. (Original) The method of claim 27, wherein said an immunological detection method is selected from the group consisting of a radio-immunoassay (RIA), an enzyme linked immunosorbent assay (ELISA), a western blot, an immunohistochemical analysis, and fluorescence activated cell sorting (FACS).
 - 29. (Cancelled)